

Supplemental legends:

Figure S1. Gene ontology (GO) analyses of the DEGs between the monokaryon strain 183 and transformant T7.

Figure S2. Kyoto Encyclopedia of Genes and Genomes (KEGG) analyses of the DEGs between the monokaryon strain 183 and transformant T7.

Table S1. Genome-wide gene expression levels of the samples.

Table S2. Differentially expressed genes (DEGs) among the samples.

Table S3. Primers used for qRT-PCR.

Figure S1

GO Enrichment

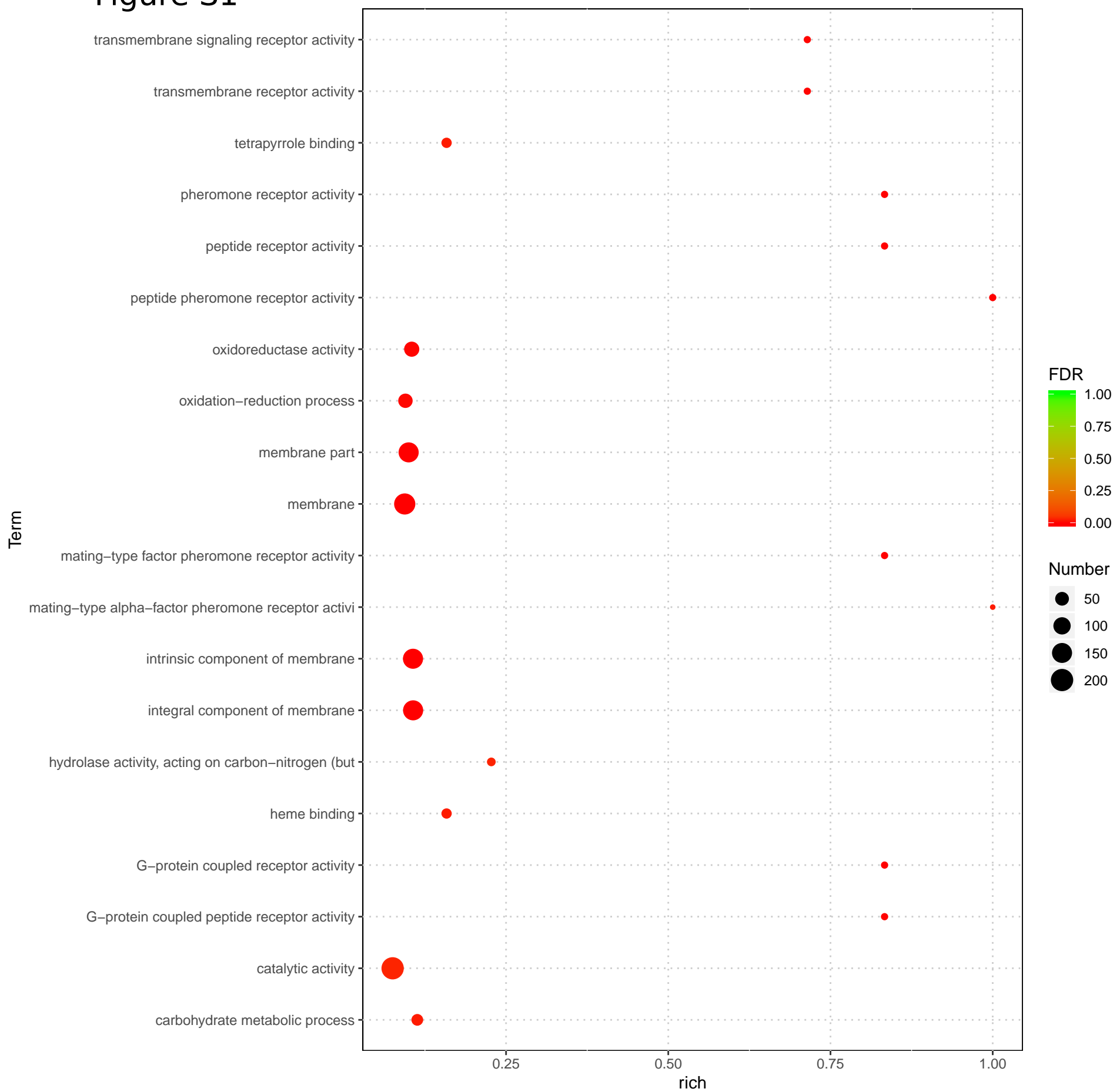


Figure S2

